

AMENDMENTS TO THE SPECIFICATION

Please enter into the specification the attached paper copy of the amended Sequence Listing, which includes 17 pages and provides sequences identified as SEQ ID NOs: 1-17.

Please replace the paragraph on page 18, line 7 to page 19, line 10, of the specification with the following paragraph:

The term "TACIs" when used herein refers to polypeptides comprising the amino acid sequence of residues 1 to ~~246~~ 247 of Figure 5B, or fragments or variants thereof, and which comprise a single cysteine rich domain. Optionally, such TACIs polypeptides comprise the contiguous sequence of residues 1 to ~~246~~ 247 of Figure 5B. Optionally, such TACIs polypeptides are encoded by the nucleic acid molecules comprising the coding polynucleotide sequence shown in Figure 5A. The TACIs polypeptides of the invention may be isolated from a variety of sources, such as from human tissue types or from another source, or prepared by recombinant and/or synthetic methods. A "native sequence" TACIs polypeptide comprises a polypeptide derived from nature. Such native sequence TACIs polypeptides can be isolated from nature or can be produced by recombinant and/or synthetic means. A TACIs polypeptide may comprise a fragment or variant of the polypeptide shown in Figure 5B and having at least about 80% amino acid sequence identity with the sequence shown in Figure 5B, more preferably at least about 81% amino acid sequence identity, more preferably at least about 82% amino acid sequence identity, more preferably at least about 83% amino acid sequence identity, more preferably at least about 84% amino acid sequence identity, more preferably at least about 85% amino acid sequence identity, more preferably at least about 86% amino acid sequence identity, more preferably at least about 87% amino acid sequence identity, more preferably at least about 88% amino acid sequence identity, more preferably at least about 89% amino acid sequence identity, more preferably at least about 90% amino acid sequence identity, more preferably at least about 91% amino acid sequence identity, more preferably at least about 92% amino acid sequence identity, more preferably at least about 93% amino acid sequence identity, more preferably at least about 94% amino acid sequence identity, more preferably at least about 95% amino acid sequence identity, more preferably at least about 96% amino acid sequence identity, more preferably at least about 97% amino acid sequence identity, more preferably at least about 98% amino acid sequence identity and yet more preferably at least about 99% amino acid sequence identity with a TACIs polypeptide encoded by an encoding nucleic acid sequence shown in Figure 5A or a specified fragment thereof. Such variant polypeptides include, for instance, polypeptides wherein one or more amino acid residues are added, or deleted, at

the N- and/or C-terminus, as well as within one or more internal domains, of the amino acid sequence shown in Figure 5B.